## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	_/0/801.487A
Source:	1FW/b
Date Processed by STIC:	6/6/06
	0//

## ENTERED



IFW16

RAW SEQUENCE LISTING DATE: 06/06/2006
PATENT APPLICATION: US/10/801,487A TIME: 08:19:57

Input Set : A:\00281FUS.txt

```
4 <110> APPLICANT: Yan, Riqiang
        Tomasselli, Alfredo G.
6
        Gurney, Mark E.
7
        Emmons, Thomas L.
        Bienkowski, Mike J.
8
        Heinrikson, Robert L.
11 <120> TITLE OF INVENTION: SUBSTRATES AND ASSAYS FOR BETA-SECRETASE ACTIVITY
13 <130> FILE REFERENCE: 29915/00281FUS
15 <140> CURRENT APPLICATION NUMBER: 10/801,487A
16 <141> CURRENT FILING DATE: 2004-03-16
18 <150> PRIOR APPLICATION NUMBER: 09/908,943
19 <151> PRIOR FILING DATE: 2001-07-19
21 <150> PRIOR APPLICATION NUMBER: 60/219,795
22 <151> PRIOR FILING DATE: 2000-07-19
24 <160> NUMBER OF SEQ ID NOS: 199
26 <170> SOFTWARE: PatentIn Ver. 2.0
28 <210> SEQ ID NO: 1
29 <211> LENGTH: 2070
30 <212> TYPE: DNA
31 <213> ORGANISM: Homo sapiens
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37 gtggagatgg tggacaacct gaggggcaag tcggggcagg gctactacgt ggagatgacc 240
38 gtgggcagcc ccccgcagac gctcaacatc ctggtggata caggcagcag taactttgca 300
39 gtgggtgctg cccccaccc cttcctgcat cgctactacc agaggcagct gtccagcaca 360
40 taccgggacc tccggaaggg tgtgtatgtg ccctacaccc agggcaagtg ggaaggggag 420
41 ctgggcaccg acctggtaag catcccccat ggccccaacg tcactgtgcg tgccaacatt 480
42 gctgccatca ctgaatcaga caagttette atcaacgget ccaactggga aggcateetg 540
43 gggctggcct atgctgagat tgccaggcct gacgactccc tggagccttt ctttgactct 600
44 ctqqtaaagc agacccacgt tcccaacctc ttctccctgc acctttgtgg tgctggcttc 660
45 cccctcaacc agtctgaagt gctggcctct gtcggaggga gcatgatcat tggaggtatc 720
46 gaccactcgc tgtacacagg cagtctctgg tatacaccca tccggcggga gtggtattat 780
47 gaggtcatca ttgtgcgggt ggagatcaat ggacaggatc tgaaaatgga ctgcaaggag 840
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52 atccttccgc agcaatacct gcggccagtg gaagatgtgg ccacgtccca agacgactgt 1140
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Input Set : A:\00281FUS.txt

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58 cgctgcctcc gctgcctgcg ccagcagcat gatgactttg ctgatgacat ctccctgctg 1500
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60 ctttggtcac aagtaggaga cacagatggc acctgtggcc agagcacctc aggaccctcc 1620
61 ccacccacca aatgcctctg ccttgatgga gaaggaaaag gctggcaagg tgggttccag 1680
62 ggactgtacc tgtaggaaac agaaaagaga agaaagaagc actctgctgg cgggaatact 1740
63 cttggtcacc tcaaatttaa gtcgggaaat tctgctgctt gaaacttcag ccctgaacct 1800
65 gtactggcat cacacgcagg ttaccttggc gtgtgtccct gtggtaccct ggcagagaag 1920
66 agaccaagct tgtttccctg ctggccaaag tcagtaggag aggatgcaca gtttgctatt 1980
67 tgctttagag acagggactg tataaacaag cctaacattg gtgcaaagat tgcctcttga 2040
68 attaaaaaaa aaaaaaaaaa aaaaaaaaaa
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72 <212> TYPE: PRT
73 <213> ORGANISM: Homo sapiens
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79 Leu Pro Ala His Gly Thr Gln His Gly Ile Arg Leu Pro Leu Arg Ser
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82 Gly Leu Gly Gly Ala Pro Leu Gly Leu Arg Leu Pro Arg Glu Thr Asp
           35
85 Glu Glu Pro Glu Glu Pro Gly Arg Arg Gly Ser Phe Val Glu Met Val
88 Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu Met Thr
                       70
                                           75
91 Val Gly Ser Pro Pro Gln Thr Leu Asn Ile Leu Val Asp Thr Gly Ser
                                       90
94 Ser Asn Phe Ala Val Gly Ala Ala Pro His Pro Phe Leu His Arg Tyr
              100
                                  105
97 Tyr Gln Arg Gln Leu Ser Ser Thr Tyr Arg Asp Leu Arg Lys Gly Val
                              120
          115
100 Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu Gly Glu Leu Gly Thr Asp
                                               140
                           135
       130
103 Leu Val Ser Ile Pro His Gly Pro Asn Val Thr Val Arg Ala Asn Ile
106 Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe Ile Asn Gly Ser Asn Trp
109 Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg Pro Asp Asp
               180
                                   185
110
112 Ser Leu Glu Pro Phe Phe Asp Ser Leu Val Lys Gln Thr His Val Pro
                               200
115 Asn Leu Phe Ser Leu His Leu Cys Gly Ala Gly Phe Pro Leu Asn Gln
                           215
       210
118 Ser Glu Val Leu Ala Ser Val Gly Gly Ser Met Ile Ile Gly Gly Ile
                       230 ·
                                           235
121 Asp His Ser Leu Tyr Thr Gly Ser Leu Trp Tyr Thr Pro Ile Arg Arg
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Input Set : A:\00281FUS.txt

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124 Glu Trp Tyr Tyr Glu Val Ile Ile Val Arg Val Glu Ile Asn Gly Gln
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                                   265
127 Asp Leu Lys Met Asp Cys Lys Glu Tyr Asn Tyr Asp Lys Ser Ile Val
                               280
           275
130 Asp Ser Gly Thr Thr Asn Leu Arg Leu Pro Lys Lys Val Phe Glu Ala
                           295
                                               300
133 Ala Val Lys Ser Ile Lys Ala Ala Ser Ser Thr Glu Lys Phe Pro Asp
                                           315
                       310
136 Gly Phe Trp Leu Gly Glu Gln Leu Val Cys Trp Gln Ala Gly Thr Thr
                                       330
                   325
137
139 Pro Trp Asn Ile Phe Pro Val Ile Ser Leu Tyr Leu Met Gly Glu Val
                                                       350
               340
                                   345
142 Thr Asn Gln Ser Phe Arg Ile Thr Ile Leu Pro Gln Gln Tyr Leu Arg
                               360
143
           355
145 Pro Val Glu Asp Val Ala Thr Ser Gln Asp Asp Cys Tyr Lys Phe Ala
                                               380
                           375
       370
148 Ile Ser Gln Ser Ser Thr Gly Thr Val Met Gly Ala Val Ile Met Glu
                                           395
                       390
149 385
151 Gly Phe Tyr Val Val Phe Asp Arg Ala Arg Lys Arg Ile Gly Phe Ala
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154 Val Ser Ala Cys His Val His Asp Glu Phe Arg Thr Ala Ala Val Glu
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                420
155
157 Gly Pro Phe Val Thr Leu Asp Met Glu Asp Cys Gly Tyr Asn Ile Pro
                                                   445
            435
                               440
158
160 Gln Thr Asp Glu Ser Thr Leu Met Thr Ile Ala Tyr Val Met Ala Ala
                                               460
                            455
163 Ile Cys Ala Leu Phe Met Leu Pro Leu Cys Leu Met Val Cys Gln Trp
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                        470
166 Arg Cys Leu Arg Cys Leu Arg Gln Gln His Asp Asp Phe Ala Asp Asp
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169 Ile Ser Leu Leu Lys
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174 <211> LENGTH: 1977
175 <212> TYPE: DNA
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182 gtggagatgg tggacaacct gaggggcaag tcggggcagg gctactacgt ggagatgacc 240
183 gtgggcagcc ccccgcagac gctcaacatc ctggtggata caggcagcag taactttgca 300
184 gtgggtgctg cccccaccc cttcctgcat cgctactacc agaggcagct gtccagcaca 360
185 taccgggacc tccggaaggg tgtgtatgtg ccctacaccc agggcaagtg ggaaggggag 420
186 ctgggcaccg acctggtaag catcccccat ggccccaacg tcactgtgcg tgccaacatt 480
187 gctgccatca ctgaatcaga caagttcttc atcaacggct ccaactggga aggcatcctg 540
188 gggctggcct atgctgagat tgccaggctt tgtggtgctg gcttccccct caaccagtct 600
189 gaagtgetgg cetetgtegg agggageatg atcattggag gtategacea etegetgtae 660
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Input Set : A:\00281FUS.txt

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191 cgggtggaga tcaatggaca ggatctgaaa atggactgca aggagtacaa ctatgacaag 780
192 agcattgtgg acagtggcac caccaacctt cgtttgccca agaaagtgtt tgaagctgca 840
193 gtcaaatcca tcaaggcagc ctcctccacg gagaagttcc ctgatggttt ctggctagga 900
194 gagcagctgg tgtgctggca agcaggcacc accccttgga acattttccc agtcatctca 960
195 ctctacctaa tgggtgaggt taccaaccag tccttccgca tcaccatcct tccgcagcaa 1020
196 tacctgcggc cagtggaaga tgtggccacg-tcccaagacg actgttacaa gtttgccatc 1080
197 tcacagtcat ccacgggcac tgttatggga gctgttatca tggagggctt ctacgttgtc 1140
198 tttgatcggg cccgaaaacg aattggcttt gctgtcagcg cttgccatgt gcacgatgag 1200
199 ttcaggacgg cagcggtgga aggccctttt gtcaccttgg acatggaaga ctgtggctac 1260
200 aacattccac agacagatga gtcaaccctc atgaccatag cctatgtcat ggctgccatc 1320
201 tgcgccctct tcatgctgcc actctgcctc atggtgtgtc agtggcgctg cctccgctgc 1380
202 ctgcgccagc agcatgatga ctttgctgat gacatctccc tgctgaagtg aggaggccca 1440
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204 ggagacacag atggcacctg tggccagagc acctcaggac cctccccacc caccaaatgc 1560
205 ctctgccttg atggagaagg aaaaggctgg caaggtgggt tccagggact gtacctgtag 1620
206 gaaacagaaa agagaagaaa gaagcactct gctggcggga atactcttgg tcacctcaaa 1680
207 tttaagtcgg gaaattctgc tgcttgaaac ttcagccctg aacctttgtc caccattcct 1740
208 ttaaattctc caacccaaag tattcttctt ttcttagttt cagaagtact ggcatcacac 1800
209 gcaggttacc ttggcgtgtg tccctgtggt accctggcag agaagagacc aagcttgttt 1860
210 ccctgctggc caaagtcagt aggagaggat gcacagtttg ctatttgctt tagagacagg 1920
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213 <210> SEQ ID NO: 4
214 <211> LENGTH: 476
215 <212> TYPE: PRT
216 <213> ORGANISM: Homo sapiens
218 <400> SEQUENCE: 4
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225 Gly Leu Gly Gly Ala Pro Leu Gly Leu Arg Leu Pro Arg Glu Thr Asp
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                                 40
228 Glu Glu Pro Glu Glu Pro Gly Arg Arg Gly Ser Phe Val Glu Met Val
                             55
231 Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu Met Thr
232 65
                         70
234 Val Gly Ser Pro Pro Gln Thr Leu Asn Ile Leu Val Asp Thr Gly Ser
235
                                          90
237 Ser Asn Phe Ala Val Gly Ala Ala Pro His Pro Phe Leu His Arg Tyr
                                    105
238
                100
240 Tyr Gln Arg Gln Leu Ser Ser Thr Tyr Arg Asp Leu Arg Lys Gly Val
                                120
                                                     125
            115
243 Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu Gly Glu Leu Gly Thr Asp
                                                 140
                            135
246 Leu Val Ser Ile Pro His Gly Pro Asn Val Thr Val Arg Ala Asn Ile
                                            155
                        150
249 Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe Ile Asn Gly Ser Asn Trp
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250
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Input Set : A:\00281FUS.txt

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252 Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg Leu Cys Gly
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               180
255 Ala Gly Phe Pro Leu Asn Gln Ser Glu Val Leu Ala Ser Val Gly Gly
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          195
258 Ser Met Ile Ile Gly Gly Ile Asp His Ser Leu Tyr Thr Gly Ser Leu
                                                220
                           215
261 Trp Tyr Thr Pro Ile Arg Arg Glu Trp Tyr Tyr Glu Val Ile Ile Val
                                            235
                       230
264 Arg Val Glu Ile Asn Gly Gln Asp Leu Lys Met Asp Cys Lys Glu Tyr
                    245
267 Asn Tyr Asp Lys Ser Ile Val Asp Ser Gly Thr Thr Asn Leu Arg Leu
                                    265
              260
270 Pro Lys Lys Val Phe Glu Ala Ala Val Lys Ser Ile Lys Ala Ala Ser
                                280
          275
273 Ser Thr Glu Lys Phe Pro Asp Gly Phe Trp Leu Gly Glu Gln Leu Val
                            295
276 Cys Trp Gln Ala Gly Thr Thr Pro Trp Asn Ile Phe Pro Val Ile Ser
                        310
                                            315
279 Leu Tyr Leu Met Gly Glu Val Thr Asn Gln Ser Phe Arg Ile Thr Ile
                                        330
                    325
283 Leu Pro Gln Gln Tyr Leu Arg Pro Val Glu Asp Val Ala Thr Ser Gln
                                    345
                340
286 Asp Asp Cys Tyr Lys Phe Ala Ile Ser Gln Ser Ser Thr Gly Thr Val
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289 Met Gly Ala Val Ile Met Glu Gly Phe Tyr Val Val Phe Asp Arg Ala
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292 Arg Lys Arg Ile Gly Phe Ala Val Ser Ala Cys His Val His Asp Glu
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                        390
295 Phe Arg Thr Ala Ala Val Glu Gly Pro Phe Val Thr Leu Asp Met Glu
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298 Asp Cys Gly Tyr Asn Ile Pro Gln Thr Asp Glu Ser Thr Leu Met Thr
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               420
301 Ile Ala Tyr Val Met Ala Ala Ile Cys Ala Leu Phe Met Leu Pro Leu
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                                440
302 435
304 Cys Leu Met Val Cys Gln Trp Arg Cys Leu Arg Cys Leu Arg Gln Gln
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307 His Asp Asp Phe Ala Asp Asp Ile Ser Leu Leu Lys
308 465
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312 <211> LENGTH: 14
313 <212> TYPE: PRT
314 <213> ORGANISM: Artificial Sequence
316 <220> FEATURE:
317 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic
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320 <400> SEQUENCE: 5
321 Lys Val Glu Ala Asn Tyr Glu Val Glu Gly Glu Arg Lys Lys
325 <210> SEO ID NO: 6
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Input Set : A:\00281FUS.txt

Output Set: N:\CRF4\06062006\J801487A.raw

## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:13; Xaa Pos. 7
Seq#:15; Xaa Pos. 4,7
Seq#:16; Xaa Pos. 1,4,5,6,7
Seq#:17; Xaa Pos. 1,2,4,5,6,7
Seq#:18; Xaa Pos. 1,2,4,5,6,7
Seq#:21; Xaa Pos. 5
Seg#:27; Xaa Pos. 7,19
Seg#:28; Xaa Pos. 6,7,11,20
Seg#:41; Xaa Pos. 9
Seq#:49; Xaa Pos. 1
Seq#:50; Xaa Pos. 2
Seq#:51; Xaa Pos. 3
Seq#:52; Xaa Pos. 4
Seq#:53; Xaa Pos. 5
Seq#:54; Xaa Pos. 6
Seq#:55; Xaa Pos. 7
Seq#:56; Xaa Pos. 8
Seq#:57; Xaa Pos. 1
Seq#:58; Xaa Pos. 2
Seq#:59; Xaa Pos. 3
Seq#:60; Xaa Pos. 4
Seq#:61; Xaa Pos. 5
Seq#:62; Xaa Pos. 6
Seq#:63; Xaa Pos. 7
Seq#:64; Xaa Pos. 8
Seq#:65; Xaa Pos. 1
Seq#:66; Xaa Pos. 2
Seq#:67; Xaa Pos. 3
Seq#:68; Xaa Pos. 4
Seq#:69; Xaa Pos. 5
Seq#:70; Xaa Pos. 6
Seq#:71; Xaa Pos. 7
Seq#:72; Xaa Pos. 8
Seq#:73; Xaa Pos. 1
Seq#:74; Xaa Pos. 2
Seq#:75; Xaa Pos. 3
Seq#:76; Xaa Pos. 4
Seq#:77; Xaa Pos. 7
Seq#:78; Xaa Pos. 8
Seq#:79; Xaa Pos. 8
Seq#:80; Xaa Pos. 9
Seq#:81; Xaa Pos. 1,7
Seq#:82; Xaa Pos. 2,7
Seq#:83; Xaa Pos. 3,7
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/801,487A

DATE: 06/06/2006 TIME: 08:19:58

Input Set : A:\00281FUS.txt

Output Set: N:\CRF4\06062006\J801487A.raw

Seq#:84; Xaa Pos. 4,7 Seq#:85; Xaa Pos. 5,7 Seq#:86; Xaa Pos. 6,7 Seq#:87; Xaa Pos. 7 Seq#:88; Xaa Pos. 7,8 Seq#:89; Xaa Pos. 1 Seq#:90; Xaa Pos. 1,2

## VERIFICATION SUMMARY

PATENT APPLICATION: US/10/801,487A

DATE: 06/06/2006 TIME: 08:19:58

Input Set : A:\00281FUS.txt

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L:476 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:500 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:524 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:595 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:695 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
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L:731 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
M:341 Repeated in SeqNo=28
L:928 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:0
L:1045 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49 after pos.:0
L:1064 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 after pos.:0
L:1083 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 after pos.:0
L:1102 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52 after pos.:0
L:1121 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53 after pos.:0
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L:1178 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56 after pos.:0
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L:1683 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82 after pos.:0
L:1707 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83 after pos.:0
L:1731 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84 after pos.:0
L:1755 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:85 after pos.:0
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VERIFICATION SUMMARY

-

DATE: 06/06/2006

PATENT APPLICATION: US/10/801,487A

TIME: 08:19:58

Input Set : A:\00281FUS.txt

Output Set: N:\CRF4\06062006\J801487A.raw

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